

INFORMATION DISCLOSURE STATEMENT	Atty. Docket No.: 265.0041 0101	Serial No.: 10/580,979
	Applicant(s): Lemon et al.	Confirmation No.: 9290
	371(c) Filing Date: April 9, 2007	Group: 1648
	Information Disclosure Statement filed: December 30, 2009	

U.S. PATENT DOCUMENTS

Examiner Initial	Copy Enclosed	Document Number	Date	Name	Class	Subclass	Filing Date If Appropriate
		None					

U.S. PATENT APPLICATIONS BY SERIAL NUMBER

Examiner Initial	Copy Enclosed	Document Number	Filing Date	Name	Class	Subclass
		None				

FOREIGN PATENT DOCUMENTS

Examiner Initial	Copy Enclosed	Document Number	Date	Country	Class	Subclass	Translation	
							Yes	No
	X	WO 2004/055216 A2, A3	07/01/04	WIPO				

OTHER DOCUMENTS (Including Authors, Title, Date, Pertinent Papers, etc.)

Examiner Initial	Copy Enclosed	Document Description
	X	Adams et al., "Complete Coding Sequence of Hepatitis C Virus Genotype 6a," <i>Biochemical and Biophysical Research Communications</i> , 1997; 234:393-396.
	X	Bressanelli et al., "Crystal structure of the RNA-dependent RNA polymerase of hepatitis C virus," <i>PNAS</i> , 1999 November 9; 96(23):13034-13039.
	X	Bressanelli et al., "Structural Analysis of the Hepatitis C Virus RNA Polymerase in Complex with Ribonucleotides," <i>Journal of Virology</i> , 2002 April; 76(7):3482-3492.
	X	Chamberlain et al., "Complete nucleotide sequence of a type 4 hepatitis C virus variant, the predominant genotype in the Middle East," <i>Journal of General Virology</i> , 1997; 78:1341-1347.

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<p>*Examiner: Initial if citation considered, whether or not citation is in conformance with MPEP 609; Draw line through citation if not in conformance and not considered. Include copy of this form with next communication to applicant.</p>	

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Examiner Initial	Copy Enclosed	Document Description
	X	Cheney et al., "Mutations in NS5B Polymerase of Hepatitis C Virus: Impacts on <i>in Vitro</i> Enzymatic Activity and Viral RNA Replication in the Subgenomic Replicon Cell Culture," <i>Virology</i> , 2002; 297:298-306.
	X	Love et al., "The Crystal Structure of Hepatitis C Virus NS3 Proteinase Reveals a Trypsin-like Fold and a Structural Zinc Binding Site," <i>Cell</i> , 1996 October 18; 87:331-342.
	X	Love et al., "Crystallographic Identification of a Noncompetitive Inhibitor Binding Site on the Hepatitis C Virus NS5B RNA Polymerase Enzyme," <i>Journal of Virology</i> , 2003 July; 77(13):7575-7581.
	X	Marzio et al., "HIV-1 Tat transactivator recruits p300 and CREB-binding protein histone acetyltransferases to the viral promoter," <i>Proc. Natl. Acad. Sci. USA</i> , 1998 November; 95:13519-13524.
	X	Simmonds et al., "A Proposed System for the Nomenclature of Hepatitis C Viral Genotypes," <i>Hepatology</i> , 1994; 19:1321-1324.
	X	Simmonds et al., "Evolutionary analysis of variants of hepatitis C virus found in South-East Asia: comparison with classifications based upon sequence similarity," <i>Journal of General Virology</i> , 1996; 77:3013-3024.
	X	Simmonds et al., "Consensus Proposals for a Unified System of Nomenclature of Hepatitis C Virus Genotypes," <i>Hepatology</i> , 2005; 42(4):962-973.
	X	Smith et al., "Characteristics of Nucleotide Substitution in the Hepatitis C Virus Genome: Constraints on Sequence Change in Coding Regions at Both Ends of the Genome," <i>J. Mol. Evol.</i> , 1997; 45:238-246.
	X	Yan et al., "Complex of NS3 protease and NS4A peptide of BK strain hepatitis C virus: A 2.2 Å resolution structure in a hexagonal crystal form," <i>Protein Science</i> , 1998; 7:837-847.

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